

# HSL mRNA sequence

>gi|21328445|ref|NM\_005357.2| Homo sapiens lipase, hormone-sensitive (LIPE), mRNA

CTTCTTGTAAGAGAGTGCTAGGCACATAGTCCCCTCCTATTCTAATCTCCCAACAAAGAAAGAGGCAC  
AGAGTTCATTACTTAGTGGGGCCAGCTGTGATCGGCCAACTGCCAGCTGCCTTAAAAAGGAAGACCAGT  
GATGCTAGGATGGAGTGAACCCAAGAGGAAGTGCCATCATGAGGAATCAATGAGAGATCTGTGAAGAGA  
GAGGGCTGGGTGGGAGCCCAAGAGGATAGAACCTGGAAGATCAATATCTCCCGTGAGGGAAATAACAATG  
GAGCCAGGTTCTAAGTCAAGTGTCTAGGTCAGACTGGCAACCTGAACACACACCAGAGGCCTATAACCCCGC  
TAGAGCCTGGGCCAGAAAAGACACCCATAGCCCAGCCAGAATCGAAGACTCTGCAGGGATCCAATACCCA  
ACAGAAGCCTGCTTCAAACCAAAAGACCCCTCACCCAGCAGGAGACCCCTGCACAACATGATGCTGAATCC  
CAGAAGGAACCTAGAGCCCAACAAAAATCTGTTTCAACAAGAGGAATTTCTTGCCCCACAGAAGCCCGCAC  
CACAGCAATCACCTTACATCCAAGGGTGCTGCTCACTCAACAGGAAGTGCTCCCAGCAGGGACCTGG  
GCTAGGAAAAGAATCTATAACTCAACAGGAGCCAGCATTGAGACAAAGACATGTAGCCCAGCCAGGGCCT  
GGCCAGGAGAGCCACCTCCAGCTCAACAAGAAAGCTGAATCAACACCTGCGGCCAGGCTAAACCTGGAG  
CCAAAAGGGAGCCATCTGCCCGACTGAATCTACGTCCCAAGAGACACCTGAACAGTCAGACAAGCAAAC  
AACGCCAGTCCAGGGGAGCCAAATCCAAGCAGGGATCTTTGACAGAGCTGGGATTTCTAACAAAACCTTCAG  
GAECTATCCATACAGCGATACGCCCTAGAGTGGAAAGGCACCTTTCTGAGTGGGTACAGATTCTGAGTCAG  
AATCAGATGTGGGATCATCTTCAGACACAGATTCTCCAGCCACGATGGGTGGAATGGTGGCCCAGGGAGT  
GAAGCTAGGCTCAAAGAAAAATCTGGTTATAAAGATGATGTCAGGATAACAGTGGGACGTCGCCACATGAG  
AAAACCAAGTCTCGGAATCAGACACTACAGGATTAAGAGTATCAGCCTCAAGGCTCATCCACAACATGGACCTGC  
GCACAATGACACAGTCGGTGGTACTCTGGCCGAGGACAACATAGCCTTCTTCTCGAGCCAGGGTCTCTGG  
GGAAACGGCCAGCGGCTGTCAAGCGTTTTTTCGGGTGTACGGGAGCAGGGCGCTGGGGCTGGAGCCCGCC  
CTCGGCCCGCTGTGGGTGTGAGCTACGCCACCGCTGTGGTTACCAATCGGCCGGGGTACTCTCTTTGAGG  
GCGACGAGGGGCTCACCGCCGACTTCTCCGGGAGTATGTCACGCTGCATAAGGGATGCTTCTATGGCCG  
CTGCCCTGGGCTTCCAGTTCACGCCTGCCATCCGGCCATTCTGCAGACCATCTCCATTGGGCTGGTGTCC  
TTGGGGAGCACTACAAAACGCAACGAGACAGGCTCAGTGTGGCCGCCAGCTCTCTTCCAGCCGGCC  
GCTTTGCCATCGACCCCGAGCTGCGTGGGGCTGAGTTTGAGCGGATCACACAGAACCTGGACGTGCACTT  
CTGGAAGCCTTCTGGAACATCACCGAGATGGAAGTGCATATCGTCTCTGGCCAACTGGCATCGGCCACC  
GTGAGGGTAAGCCGCTGCTCACCTGCCACCCGAAGCCTTTGAGATGCCACTGACTGCGCACCCACGC  
TCACGGTCACCATCTCACCCCCACTGGCCACACAGGCCCTGGGCCGCTCCTCGTCAGGCTCATCTCCTA  
TGACCTGCGTGAAGGACAGGACAGTGAAGGACTCAGCAGCCTGATAAAGTCCAACGGCCAACGGAGCCGTG  
GAGCTGTGGCCGCGCCCCAGCAGGACCCCGCTCGCGGTCCTGATAGTGCACCTCCACGGCGGTGGCT  
TTGTGGCCAGACCTCCAGATCCCAAGAGCCCTACCTCAAGAGCTGGGCCAGGAGCTGGGGCCCCCAT  
CATCTCCATCGACTACTCCCTGGCCCTGAGGCCCTTCCCCCGTGCCTGGAGGAGTGTCTTTCGCC  
TACTGCTGGGCCATCAAGCACTGCGCCCTCCTTGCTCAACAGGGGAACGAATCTGCCTTGCGGGGGACA  
GTGCAGGCGGGAACTCTGCTTACCGTGCTCTTCCGGCAGCAGCCTACGGGGTGCGGGTGCCAGATGG  
CATCATGGCAGCCTACCCGGCCACAATGCTGACGCCTGCCGCCTCTCCCTCCCGCTGCTGAGCCTCATG  
GACCCCTTGCTGCCCTCAGTGTGCTCTCCAAGTGTGTACAGCCCTATGCTGGTGCAAAGACGGAGGACC  
ACTCCAACCTCAGACCAGAAAGCCCTCGGCATGATGGGGCTGGTGCGGCCGGGACACAGCCCTGCTCCTCG  
AGACTTCCGCCTGGGTGCCTCCTCATGGCTCAACTCCTTCTGGAGTTAAGTGGGCGCAAGTCCCAGAAG  
ATGTCCGAGCCCATAGCAGAGCCGATGCGCCGCAAGTGTGTCTGAAGCAGCACTGGCCCAGCCCCAGGGCC  
CACTGGGCACGGATTCCCTCAAGAACCTGACCCTGAGGGACTTGAGCCTGAGGGGAAACTCCGAGACGTC  
GTCGGACACCCCGAGATGTGCTGTGAGCTGAGACACTTACCCCTCCACACCCCTCCGATGTCAACTTC  
TTATTACCACCTGAGGATGACAGGGAAGAGGCTGAGGCCAAAAATGAGCTGAGCCCATGGACAGAGGCC  
TGGGCGTCCGTGCCGCTTCCCCGAGGGTTTCCACCCCGACGCTCCAGCCAGGGTGCCACACAGATGCC  
CCTCTACTCCTCACCCATAGTCAAGAACCCCTTATGTCGCCGCTGCTGGCACCCGACAGCATGCTCAAG  
AGCCTGCCACCTGTGCACATCGTGGCGTGCAGCCTGGACCCCATGCTGGACGACTCGGTCATGCTCGCGC  
GGGACTGGCAACCTGGGCCAGCCGGTGCAGCTGCGCGTGGTGGAGGACCTGCCGCAGGGCTTCTGAC  
CCTAGCGCGCTGTGCCGCGAGACGCGCCAGGCCGACAGCTGTGCGTGGAGCGCATCCGCCTCGTCTC  
ACTCCTCCCGCCGAGCCGGGGAGCGGAGCAGCCACACACACCCGCTACCCAGACGGCTGGACCTGCACGCCCTGC  
GACACTAAAAGCCTGTTGTTCCCATCTGCGCCGGCTCCGTCATGAATGCCCTCCGGGCCGGGCGGAAGG  
GGACGCGGGCTGTGCCCTACTTAAGTCGGGGTGGCAAGGGGGCGGGGGCCGAAAGCTGAGACC  
CTCGCCACGGGGAGGGGGACGCGCACACACACCCGCTACCCAGACGGCTGGACCTGCACGCCCTGC  
CTTTGCTGCTGCTGCTGCGGGCAGCCGACGGGGACTGGCCCTCCTTGACAGTGGTTTGGT TTGTTGTAATAAAAGTATTTAATTA